NUCLEOTIDE AND AA SEQUENCE

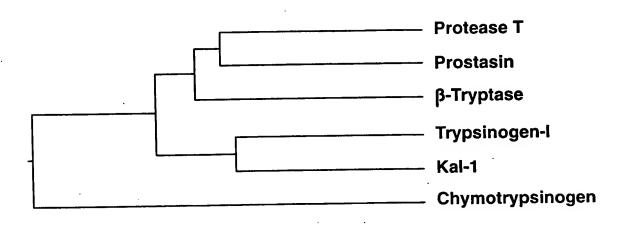
Nucleic acid sequence of Protease T (SEQ.ID.NO.:1)

GACCACGGCCTGCGCCCAGCCAGGCCTGAGGACATGAGGCGGCGGCGGCGGTGCCGC TCCTGCTGCTGTGTTTTGGGTCTCAGAGGGCCAAGGCAGCAACAGCCTGTGGTCGCC CCAGGATGCTGAACCGAATGGTGGGCGGGCAGGACACGCAGGAGGGCGAGTGGCCCTGGC AAGTCAGCATCCAGCGCAACGGAAGCCACTTCTGCGGGGGCAGCCTCATCGCGGAGCAGT GGGTCCTGACGGCTGCGCACTGCTTCCGCAACACCTCTGAGACGTCCCTGTACCAGGTCC AGGTGGAGAGCAACCCCCTGTACCAGGGCACGGCCTCCAGCGCTGACGTGGCCCTGGTGG AAGACCTCCTGCCCGAACCGCGGATCCTGCAGAAACTCGCTGTGCCCATCATCGACACAC CCAAGIGCAACCTGCTCTACAGCAAAGACACCGAGTTTGGCTACCAACCCAAAACCATCA AGAATGACATGCTGTGCGCCGGCTTCGAGGAGGGCCAAGAAGGATGCCTGCAAGGGCGACT CGGGGGGCCCCTGGTGTGCCTCGTGGGTCAGTCGTGGCTGCAGGCGGGGGTGATCAGCT GGGGTGAGGGCTGTGCCCGCCAGAACCGCCCAGGTGTCTACATCCGTGTCACCGCCCACC AGAAGTGAGACCCCCGGGGCCAGGAGCCCCTTGAGCAGAGCTCTGCACCCAGCCTGCCCG CCCACACCATCCTGCTGCTCCCCAGCGCTGCTGTTGCACCTGTGAGCCCCACCAGACT CATTIGTAAATAGCGCTCCTTCCTCCCCCTCTCAAATACCCTTATTTTATTTTATGTTTCTC CCAATAAAAACCCAGCCTGTGTGCCAGCTG

>Amino Acid Sequence of Protease T (SEQ.ID.NO.:7)

MRRPAAVPILLILCFGSQRAKAATACGRPRMLNRMVGGQDTQEGEWPWQVSIQRNGSHFC GGSLIAEQWVLTAAHCFRNTSETSLYQVLLGARQLVQPGPHAMYARVRQVESNPLYQGTA SSADVALVELEAPVPFTNYILPVCLPDPSVIFETGMVCWVTGWGSPSEEDLLPEPRILQK LAVPIIDTPKCNLLYSKDTEFGYQPKTIKNDMLCAGFEEGKKDACKGDSGGPLVCLVGQS WLQAGVISWGEGCARQNRPGVYIRVTAHHNWIHRIIPKLQFQPARLGGQK

FIG. 2



Prot.

ال 10.

SKELETAL MUSCLE LYMPH NODE MONOCYTE (PBM) OSTEOSARCOMA PANCREAS PITUITARY **PLACENTA PROSTATE** RETINA OVARY

Prof. T

HIPPOCAMPUS
HYPOTHALAMUS
SUBSTANTIA NIGRA
SUBTHALMIC NUCLEUS BREAST CARCINOMA CEREBELLUM CEREBRAL CORTEX **BCELL LYMPHOMA BONE MARROW** CAUDATE NUCL **THALAMUS AMYGDALA**

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SMALL INTESTINE **VEG CONTROL NEG CONTROL** NEG CONTROL SPINAL CORD SPLEEN J937+PMA STOMACH **JTERUS THYMUS ESTIS** 1937

JURKAT+PMA+PHA-M KERATINOCYTE COLON ENDOTHELIAL FETAL BRAIN FIBROBLAST HEART **EUKOCYTE EPIDERMIS** -AT CELL KIDNEY JURKAT

FUSION PROTEIN

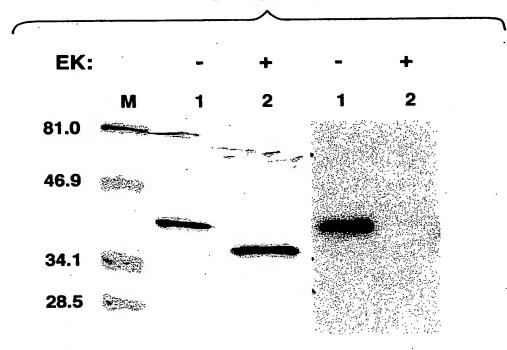
Nucleic Acid sequence of PFEK-PROTT-HIS fusion protein (SEQ.ID.NO.:8).

GAATTCACCACCATGGACAGCAAAGGTTCGTCGCAGAAATCCCGCCTGCTCCTGCTGCTG GTGGTGTCAAATCTACTCTTGTGCCAGGGTGTGGTCTCCGACTACAAGGACGACGACGAC GTGGACGCGCCCCTTTGCTGCCCCCTTTGATGATGATGACAAGATCGTTGGGGGCTAT GCTCTAGAGGAGGGCGAGTGGCCCTGGCAAGTCAGCATCCAGCGCAACGGAAGCCACTTC TGCGGGGCAGCTCATCGCGGAGCAGTGGGTCCTGACGCTGCGCACTGCTTCCGCAAC ACCTCTGAGACGTCCTGTACCAGGTCCTGCTGGGGGGCAAGGCAGCTAGTGCAGCCGGGA CCACACGCTATGTATGCCCGGGTGAGGCAGGTGGAGAGCAACCCCCTGTACCAGGGCACG GCCTCCAGCGCTGACGTGGCCCTGGTGGAGCTGGAGCACCAGTGCCCTTCACCAATTAC ATCCTCCCGTGTGCCTGACCCCTCGGTGATCTTTGAGACGGGCATGAACTGCTGG GTCACTGGCTGGGGCAGCCCCAGTGAGGAAGACCTCCTGCCCGAACCGCGGATCCTGCAG AAACTCGCTGTGCCCATCATCGACACACCCAAGTGCAACCTGCTCTACAGCAAAGACACC GAGTTTGGCTACCAACCCAAAACCATCAAGAATGACATGCTGTGCGCCGGCTTCGAGGAG GGCAAGAAGGATGCCTGCAAGGGCGACTCGGGCGCCCCCTGGTGTGCCTCGTGGGTCAG TCGTGGCTGCAGGCGGGGTGATCAGCTGGGGTGAGGGCTGTGCCCGCCAGAACCGCCCA GGTGTCTACATCCGTGTCACCGCCCACCACAACTGGATCCATCGGATCATCCCCAAACTG CAGTTCCAGCCAGCGAGGTTGGGCGGCCAGAAGTCTAGACATCACCATCACCATCACTAG CGGCCGCTTCCCTTTAGTGAGGGTTAATGCTTCGAGCAGACATGATAAGATACATTGATG AGTTTGGACAAACCACAACTAGAATGCAGTGAAAAAAATGCTTTATTTGTGAAATTTGTG ATGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAAACAAGTTGAC

Amino Acid sequence of PFEK-PROTT-HIS fusion protein (SEQ.ID.NO.:9).

MDSKGSSQKSRLLLLLVVSNLLLCQGVVSDYKDDDDVDAAALAAPFDDDDKTVGGYALEE GEWPWQVSIQRNGSHFCGGSLIAEQWVLTAAHCFRNTSETSLYQVLLGARQLVQPGPHAM YARVRQVESNPLYQGTASSADVALVELEAPVPFTNYILPVCLPDPSVIFETGMNCWVTGW GSPSEEDLLPEPRILQKLAVPIIDTPKCNLLYSKDTEFGYQPKTIKNDMLCAGFEEGKKD ACKGDSGGPLVCLVGQSWLQAGVISWGEGCARQNRPGVYIRVTAHHNWIHRIIPKLQFQP ARLGGQKSRHHHHHH





Protease: PFEK-Protease T-HIS

Protease Activity of Protease T

H-D-Pro-HHT-Arg-pNA
H-D-Lys(CBO)-Pro-Arg-pNA
H-DL-Val-Leu-Arg-pNA

